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## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/524,426  
Source: PU/10  
Date Processed by STIC: 1/30/06

***ENTERED***



PCT

## RAW SEQUENCE LISTING

DATE: 01/23/2006

PATENT APPLICATION: US/10/524,426

TIME: 09:14:26

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\01232006\J524426.raw

3 <110> APPLICANT: Li, Limin  
 4 Aghdasi, Bahman  
 6 <120> TITLE OF INVENTION: MAMMALIAN GENES INVOLVED IN RAPAMYCIN RESISTANCE  
 AND  
 7 TUMORGENESIS: RAPR7 GENES  
 9 <130> FILE REFERENCE: 70017.11USWO  
 11 <140> CURRENT APPLICATION NUMBER: US 10/524,426  
 12 <141> CURRENT FILING DATE: 2005-02-15  
 14 <150> PRIOR APPLICATION NUMBER: PCT/US2003/026073  
 15 <151> PRIOR FILING DATE: 2003-08-15  
 17 <150> PRIOR APPLICATION NUMBER: US 60/404,311  
 18 <151> PRIOR FILING DATE: 2002-08-15  
 20 <160> NUMBER OF SEQ ID NOS: 23  
 22 <170> SOFTWARE: PatentIn version 3.3  
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 26 <212> TYPE: DNA  
 27 <213> ORGANISM: Murine  
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 58 <223> OTHER INFORMATION: n is a, c, g, or t  
 60 <220> FEATURE:  
 61 <221> NAME/KEY: misc\_feature

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62 <222> LOCATION: (931)..(931)  
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122 <222> LOCATION: (1165)..(1165)

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138 &lt;223&gt; OTHER INFORMATION: n is a, c, g, or t

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141 &lt;221&gt; NAME/KEY: misc\_feature

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      155 tgcccttttg caggttctct tactgaccat cccacctgc cccacacatc ctcccctatg      180
W--> 157 caccccaact ntgagccctt cctgtctcagt aagtctgtag acttggtggg tatattggnc      240
      159 tcattgagac tgcaggccct tggagggcag gctctgacct gcagtaagat gtgtgagtga      300
W--> 161 tactcagcac acantaggtg gataaatacc cccacagtag gtgggtagtg agccctgtga      360
W--> 163 gtccactgta agncaccatc tacatgggca nagectgctt taagcgtggg ttagggacac      420
      165 aacagtctct tcagcagggc ttctggcacc atctacacaa gtccatcctc agctcttcca      480
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      169 cctttcaact gagcccttgg ctcttggagt tagccacaac ctaactactc aggtccctcc      600
      171 aacaagggga ctgtgtctgt ggctggatga ctcatgcaca ctgctccatc ccgcaatctt      660
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      175 gcccacaacc agccctacca gaacagagtg tactcagagc tccaggacaa aaatctggaa      780
      177 acagagagcc ggctctcatt tggaccgaga tctgagtgat gaaaagagca ggcagaggaa      840
      179 acagcaagtt caaagttcct gaggtgggaa tgcgcttgac acaacggaga cctgagaaga      900
W--> 181 acacagcaaa ggccgtgtta catttgtctg ngactccagc ccccaaggat ctggtcagga      960
W--> 183 cagacatngc gaggactcac ctggataatc cagagccatg gcccatacna ngmntncttc     1020
W--> 185 tttttttttt ttctctcttt tttctttttt tctttttttg nnnnnngccc caagacaggc     1080
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W--> 191 ccactggcca ggctaactaa ggttcttaac tttttaagna ttatttttct ttcttatgta     1260
      193 tgtgtatatg ggggagggga tgcacaaggg catggggggg ggggtccctgc agaagtccga     1320
      195 agaggtgcca gatccctggg agctggaatt aaagtcagtc atgaaacatc caagatggac     1380
W--> 197 actgggnaac tgaacttggg tcctctgcga gaggagtaat ggtcttaact gctgagccat     1440
      199 ctctaggccc aatgtctggt tttgttttgt tttgttttgt tttgttttgt tttgttttgt     1500
      201 tttgtatttg ggggtttttg tttgtctgtt tgggtgggtt gtttgggttt tcttgagaca     1560
      203 gggtttctct gtatagccct ggctgtcctg gaactcactc tatagactag gctggcctcg     1620

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205 aactcagaaa tctctctgcc tctgctctcc aagtgtctgag attaaaggcc cgtgccacca 1680  
 207 ctgccccgacg ccaatgtctg tattttattc atctctgcag aatctctttt gtctcctaac 1740  
 209 ggaacatcat cccagattct gggaagtaca ctgaagacaa tgggggtgggt gttgtttctc 1800  
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 217 <211> LENGTH: 2856  
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 219 <213> ORGANISM: Murine  
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 226 cacatctcca ccatcaacca cacccttcca tctttctctt catctgacac atatcttcca 180  
 228 acccttcagt catctaataa gcagacttta aaagccacgg gtccctggata tccaatggaa 240  
 230 aatgaccaaa ggaagaacac ttgctcctta gtccgacaa aaggtttcaa aggagtccac 300  
 232 ttgcatgctg aagcacttcc cacagaagga gcaccccccc cccacacctca tctgcaggat 360  
 234 tccgagatgg aagagaagag gcgaaaatat tccatcagca gcgacaactc tgataccact 420  
 236 gacggtcacg tgacatccac atcagcatca agatgttcca aactgcccgag cagcaccaag 480  
 238 tcgggctggc cccggcagaa cgagaagaag ccctcagagg ttttcgggac agacttgatc 540  
 240 acagccatga agatcccaga ttcataccag ctacgcccgg atgactacta catcctggcg 600  
 242 gacccgtggc gacaagaatg ggagaaagg gtgcaggtag ctgctggagc ggaggccatt 660  
 244 ccagagcctg tggtaggct cctcccacca ctgaaaggcc cccccacgca gatgtcccca 720  
 246 gatagcccca cacttggtga ggggtgccc atctgactggc caggaggcag ccgctacgac 780  
 248 ctggatgaga tcgatgcgta ctggttgga cttctcaact cggagctcaa ggagatggag 840  
 250 aagcccagac tggatgagct aacgttagag cgtgttctag aggagctaga gacattgtgc 900  
 252 caccagaata tggcacaggc cattgagaca caggaggggc tgggcatcga gtacgacgag 960  
 254 gacgttgtct gcgacgtgtg ccgttcccc gaaggcgagg atggcaacga gatggtcttc 1020  
 256 tgtgacaaat gcaatgtctg tgtgcaccag gcatgctacg ggatcctcaa ggtgcctacg 1080  
 258 ggcagctggc tgtgccggac ctgtgcctg ggagtcacgc ctaagtgcct gctctgcccc 1140  
 260 aagcgaggag gagccctgaa gccactaga agtgggacca agtgggtaca cgtcagctgt 1200  
 262 gccctgtgga ttcctgaggt cagcattggc tgtccagaga agatggagcc cattaccaag 1260  
 264 atctcgcata ttccggccag ccgctgggccc ctgtcctgca gcctctgcaa ggagtgcaca 1320  
 266 ggtacctgca tccagtgttc catgccttcc tgcacacag cattccacgt tacgtgcgcc 1380  
 268 tttgaccgag gcctggaaat gcggactata ttagctgaca atgacgaggt caagttcaag 1440  
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 272 gagcccagcc aggcctgtga ggatctggaa aaggtgacct tacgcaagca gcggctgcag 1560  
 274 cagctggaag aaaacttcta tgagctagtg gagccagctg aggtggctga acggctagac 1620  
 276 ctggctgagg cactggtgga cttcatctac cagtactgga agttgaagcg gagagctaata 1680  
 278 gccaaaccagc cgctgttgac gcccaagact gacgaggtgg acaacctggc ccaacaggaa 1740  
 280 caggatgtcc tctatcgacg cctgaagctt ttcaccacc tgcggcagga cctggagagg 1800  
 282 gtaaggaacc tgtgctacat ggtgacaaga cgggagagaa cgaaacacac catctgtaaa 1860  
 284 cttcaggagc agatattcca tctacagatg aaacttattg agcaagacct ttgcagagag 1920  
 286 ctttctggga ggaggtcaaa gggcaagaag aatgattcaa aaaggaaagg ccgagagggg 1980  
 288 cccaagggca gcagccctga gaagaaagag aaagtgaagg ctgggcccga gtctgtgctg 2040  
 290 gggcagctgg gtctatccac ctctgtcccc atcgacggca ctttcttcaa cagctggttg 2100  
 292 gcacagtcgg ttcagatcac agcagaggac atggccatga gcgagtggtc tttgaacagt 2160  
 294 gggcaccggg aggatcctgc tccaggtctg ctgtcagagg aattgctaca agatgaggag 2220  
 296 acgctgctca gcttcatgag ggaccctcgc ctacgacctg gtgacctgc cagaaaggcc 2280  
 298 cgaggccgca ctgcctgcc tgccaagaag aaacctccc cgctgcagga tgggcccagt 2340

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300 gcacggacca ctccagacaa gcaacccaag aaggcctggg cccaggatgg caaggggacg      2400
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304 gcagctgggg actgtccagt cccagcaaca ctggaaagcc ctccaccact ggctcccgag      2520
306 atactagaca agacagcccc catggcttcc gacttaaata tccaagtgcc tggccctaca      2580
308 gtgagcccca aacccttggg caggctccgg ccaccccgag agatgaaggt cagtcggaaa      2640
310 tctccgggtg ctagatccga tgctgggaca ggactaccgt ctgctgtggc cgagaggcca      2700
312 aaggtcagcc tgcattttga caccgaggct gacggctact tctctgatga ggagatgagc      2760
314 gattctgagg tagaggcaga ggacagtggg gtacaacgag cttccaggga ggcaggggca      2820
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320 <211> LENGTH: 951
321 <212> TYPE: PRT
322 <213> ORGANISM: Murine
324 <400> SEQUENCE: 3
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331                20                    25                    30
334 Ser Leu Asn Leu Pro Pro Leu Cys His Ile Ser Thr Ile Asn His Thr
335                35                    40                    45
338 Leu Pro Ser Phe Ser Ser Ser Asp Thr Tyr Leu Pro Thr Leu Gln Ser
339                50                    55                    60
342 Ser Asn Lys Gln Thr Leu Lys Ala Thr Gly Pro Gly Tyr Pro Met Glu
343 65                    70                    75                    80
346 Asn Asp Gln Arg Lys Asn Thr Cys Ser Leu Val Arg Gln Glu Gly Phe
347                85                    90                    95
350 Lys Gly Val Thr Leu His Ala Glu Ala Leu Pro Thr Glu Gly Ala Pro
351                100                   105                   110
354 Pro Pro Pro Pro His Leu Gln Asp Ser Glu Met Glu Glu Lys Arg Arg
355                115                   120                   125
358 Lys Tyr Ser Ile Ser Ser Asp Asn Ser Asp Thr Thr Asp Gly His Val
359                130                   135                   140
362 Thr Ser Thr Ser Ala Ser Arg Cys Ser Lys Leu Pro Ser Ser Thr Lys
363 145                   150                   155                   160
366 Ser Gly Trp Pro Arg Gln Asn Glu Lys Lys Pro Ser Glu Val Phe Arg
367                165                   170                   175
370 Thr Asp Leu Ile Thr Ala Met Lys Ile Pro Asp Ser Tyr Gln Leu Ser
371                180                   185                   190
374 Pro Asp Asp Tyr Tyr Ile Leu Ala Asp Pro Trp Arg Gln Glu Trp Glu
375                195                   200                   205
378 Lys Gly Val Gln Val Pro Ala Gly Ala Glu Ala Ile Pro Glu Pro Val
379                210                   215                   220
382 Val Arg Leu Leu Pro Pro Leu Lys Gly Pro Pro Thr Gln Met Ser Pro
383 225                   230                   235                   240
386 Asp Ser Pro Thr Leu Gly Glu Gly Ala His Pro Asp Trp Pro Gly Gly
387                245                   250                   255
390 Ser Arg Tyr Asp Leu Asp Glu Ile Asp Ala Tyr Trp Leu Glu Leu Leu
391                260                   265                   270
394 Asn Ser Glu Leu Lys Glu Met Glu Lys Pro Glu Leu Asp Glu Leu Thr

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:1; N Pos. 1061,1062,1063,1064,1065,1089,1113,1117,1137,1159,1165,1168  
Seq#:1; N Pos. 1198,1239,1387,1818

## VERIFICATION SUMMARY

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L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180  
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:300  
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:360  
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:900  
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:960  
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1020  
L:187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1080  
L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1140  
L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1200  
L:197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1380  
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1800